1	MENGTHVNG <mark>S</mark> ANG <mark>F</mark> CIKDPLNWGVAAEALTGSHLDEVKKMVGEFRKPVVK MENG <mark>N</mark> HVNGV <mark>V</mark> NE <mark>LCIKDPLNWGVAAEALTGSHLDEVKKMVA</mark> EFRKPVVK	Sunflower LsPAL1
51	LGGETLTVSQVAGI <mark>s</mark> aagdg <mark>nm</mark> vkvelseaaragvkassdwvmesmnkgt	Sunflower
51	LGGETLTVSQVAGIAAA <mark>n</mark> D <mark>S</mark> DTVKVELSEAARAGVKASSDWVMESMNKGT	LsPAL1
101	DSYGVTTGFGATSHRRTK <mark>N</mark> GGALQKELIRFLNAGIFGNGTESSHTLPHSA	Sunflower
101	DSYGVTTGFGATSHRRTK GGALQKELIRFLNAGIFGNGTE <mark>T</mark> HTLPHSA	LsPAL1
151	TRAAMIVRINTLLQGYSGIRFEILEAITKFLNNNITPCLPLRGTITASGD	Sunflower
151	TRAAMIVRINTLLQGYSGIRFEILEAITKFLN N TP LPLRGTITASGD	LsPAL1
201	LVPLSYIAGLLTGRPNSKAVGPAGEVLNAESAFA <mark>P</mark> AGVEGGFFELQPKEG	Sunflower
201	LVPLSYIAGLLTGR NSKAVGP <mark>T</mark> GEVLNAE <mark>K</mark> AFA AGVEGGFFELQPKEG	LsPAL1
251	LALVNGTAVGSGMASMVLF <mark>D</mark> ANVLALLSEVLSA I FAEVMQGKPEFTDHLT	Sunflower
251	LALVNGTAVGSGMASMVLFDANVLALLSEVLSA I FAEVMQGKPEFTDHLT	LsPAL1
301	HKLKHHPGQIEAAAIMEYILDGSDYVKAAQKVHEMDPLQKPKQDRYALRT	Sunflower
301	HKLKHHPGQIEAAAIMEYILDGSDYVKAAQKVHEMDPLQKPKQDRYALRT	LsPAL1
351	SPQWLGPQIEVIRSATKMIEREINSVNDNPLIDVSRNKALHGGNFQGTPI	Sunflower
351	SPQWLGPQIEVIRS <mark>S</mark> TKMIEREINSVNDNPLIDVSRNKALHGGNFQGTPI	LsPAL1
401	GVSMDNTRLA1AA1GKVT1AQFSELVNDFYNNGLPS <mark>H</mark> LSGGRNPSLDSGF	Sunflower
401	GVSMDNTRLA1AA1GK <mark>LMF</mark> AQFSELVNDFYNNGLPSNLSGGRNPSLD <mark>Y</mark> GF	LsPAL1
451	KGGE LAMASYCSELQFLANPVTNHVQSAEQHNQDVNSLGL I SARKTAEAV	Sunflower
451	KGGE LAMASYCSELQFLANPVTNHVQSAEQHNQDVNSLGL I SARKTAEAV	LsPAL1
501	DILKLMSSTYLVALCQSIDLRHLEENMKSTVKNTVSQVAKKVLTMGVNGE	Sunflower
501	DILKLMSSTYLVALCQSIDLRHLEENMKSTVKNTVSQVAKKVLTMGVNGE	LsPAL1
551	LHPSRFCEKDLLRVVDREYVFAYADD <mark>P</mark> CL <mark>T</mark> TYPLMQKLRQVLVDHALNNG	Sunflower
551	LHPSRFCEKDLLRVVDREYVFAY <mark>I</mark> DDVC <mark>S</mark> GTYPLMQKLRQVLVDHALNNG	LsPAL1
601	ETEKNANTSIFQKIATFEDELKA <mark>I</mark> LPKEVE <mark>S</mark> VRVAFENGT <mark>M</mark> SIPNRIKAC	Sunflower
601	ETEKN <mark>T</mark> NTSIFQKIATFE <mark>E</mark> ELK <mark>V</mark> LLPKEVEGVR <mark>I</mark> A <mark>Y</mark> EN <mark>D</mark> TLSIPNRIKAC	LsPAL1
651	RSYPLYRFVREELGG <mark>A</mark>	Sunflower
651	RSYPLYRFVREELG <mark>R</mark> GFLTGEKVTSPGEEFDRVFTAMCKGQIIDPLLECL	LsPAL1
701	GGWNGEPLPIC	LsPAL1